## **AMENDMENTS TO THE CLAIMS:**

Claims 1-39 are canceled without prejudice or disclaimer. Claims 40-47 are added. The following is the status of the claims of the above-captioned application.

Claims 1-39 (Cancelled)

Claim 40. (New.) A variant of a parent Termamyl-like α-amylase, wherein said variant has α-amylase activity and at least 80% sequence identity to said parent α-amylase and comprises a deletion at a position corresponding to position R181 or G182 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 41. (New.) The variant of claim 40, wherein the variant comprises a deletion at a position corresponding to position R181 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 42. (New.) The variant of claim 40, wherein the variant comprises a deletion at a position corresponding to position G182 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 43. (New.) The variant of claim 40, wherein the variant comprises a deletion at a position corresponding to position R181 and G182 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 44. (New.) A variant of a parent Termamyl-like  $\alpha$ -amylase, wherein said variant has  $\alpha$ -amylase activity and at least 80% sequence identity to said parent  $\alpha$ -amylase and comprises a mutation at a position corresponding to position G149 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 45. (New.) The variant of claim 44, wherein the variant comprises a mutation at a position corresponding to position G149A in the amino acid sequence shown in SEQ ID NO: 2.

Claim 46. (New.) A variant of a parent Termamyl-like  $\alpha$ -amylase, wherein said variant has  $\alpha$ -amylase activity and at least 80% sequence identity to said parent  $\alpha$ -amylase and comprises a mutation at a position corresponding to position H107 in the amino acid sequence shown in SEQ ID NO: 2.

Claim 47. (New.) The variant of claim 44, wherein the variant comprises a mutation at a position corresponding to position H107I,L,V,F,Y, or K in the amino acid sequence shown in SEQ ID NO: 2.